



SEQUENCE LISTING

#8
<110> Madsen, Soren
Vrang, Astrid
Bredmose, Lars
Ravn, Peter
Glenting, Jacob
Johnson, Mads Gronvald
Israelsen, Hans

<120> REGULATION OF PROMOTER ACTIVITY IN CELLS

<130> 54320.000010

<140> US 09/982,532

<141> 2001-10-19

<160> 10

<170> PatentIn version 3.1

<210> 1

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Cy5 labelled ISS1.F1 primer

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TECH CENTER 1600/2900

APR 15 2002

RECEIVED

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120

gttacagccc tgtatatggc gaaataaaatg aataaaaaat agcgagttaga tgagttttaa

180

aatgaaaagaa atggcaaacg taaacattga atatctaattc aatacactgg aacaaaaaaaaa

240

agtgagtgtt gtaacacgca aaaaacatag ttatcatg tatcaaggaa ttgaatcaga

300

atatatctat gtactcaaag atgggttagc gaagattagc aatattttaa gagatggtcg	360
tgaatttaat attgcattatg ttgcggagcc agactttgtt tctttattgg aagagaaaaca	420
aaacgatgga atttcagcat tatttaatgt acgaatttagt tctccaacag ccagtttta	480
caaaaattca cgcagtgatt tttggaattt ggttcgttag gatttgaatt tattcagagt	540
tgttgatgac ttttataaac gaagactagc acttaattta gaaattcttc aaaagatgac	600
aatcaatggt aagaagggag cggttgcgc ttgccttcac agttttagt atgatttcgg	660
aataagaaaa aaagatggaa ttctgattga ttttaccgtc actaatgaag atattgcagg	720
tttttgttgtt atttctacac gaaatagtgt taaccgtatt ctcatgatt taaaggatga	780
aaaagtaatt ggagttagt ataataaaat tatgattat aatcctcaat acttagaaga	840
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His Ser Tyr Ile Met Tyr Gln Gly Ile Glu Ser Glu Tyr Ile Tyr Val			
35	40	45	

Leu Lys Asp Gly Val Ala Lys Ile Ser Asn Ile Leu Arg Asp Gly Arg			
50	55	60	

Glu Phe Asn Ile Ala Tyr Val Ala Glu Pro Asp Phe Val Ser Leu Leu			
65	70	75	80

Glu Glu Lys Gln Asn Asp Gly Ile Ser Ala Leu Phe Asn Val Arg Ile			
85	90	95	

Glu Ser Pro Thr Ala Ser Phe Tyr Lys Ile Ser Arg Ser Asp Phe Trp			
100	105	110	

Asn Trp Val Arg Glu Asp Leu Asn Leu Phe Arg Val Val Asp Asp Phe			
115	120	125	

Tyr Lys Arg Arg Leu Ala Leu Asn Leu Glu Ile Leu Gln Lys Met Thr

130

135

140

Ile Asn Gly Lys Lys Gly Ala Val Cys Ala Cys Leu His Ser Leu Ile
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Asp Asp Phe Gly Ile Arg Lys Lys Asp Gly Ile Leu Ile Asp Phe Thr
165 170 175

Val Thr Asn Glu Asp Ile Ala Gly Phe Cys Gly Ile Ser Thr Arg Asn
180 185 190

Ser Val Asn Arg Ile Leu His Asp Leu Lys Asp Glu Lys Val Ile Gly
195 200 205

Val Ile Asp Asn Lys Ile Met Ile Tyr Asn Pro Gln Tyr Leu Glu Glu
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Tyr Ile Ser
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Leu Lys Asp Gly Val Ala Lys Ile Ser Asn Ile Leu Arg Asp Gly Arg
50 55 60

Glu Phe Asn Ile Ala Tyr Val Ala Glu Pro Asp Phe Val Ser Leu Leu
65 70 75 80

Glu Glu Lys Gln Asn Asp Gly Ile Ser Ala Leu Phe Asn Val Arg Ile
85 90 95

Glu Ser Pro Thr Ala Ser Phe Tyr Lys Ile Ser Arg Ser Asp Phe Trp
100 105 110

Asn Trp Val Arg Glu Asp Leu Asn Leu Phe Arg Val Val Asp Asp Phe
115 120 125

Tyr Lys Arg Arg Leu Ala Leu Asn Leu Glu Ile Leu Gln Lys Met Thr
130 135 140

Ile Asn Gly Lys Lys Gly Ala Val Cys Ala Cys Leu His Ser Leu Ile
145 150 155 160

Asp Asp Phe Gly Ile Arg Lys Lys Asp Gly Ile Leu Ile Asp Phe Thr
165 170 175

Val Thr Asn Glu Asp Ile Ala Gly Phe Cys Gly Ile Ser Thr Arg Asn
180 185 190

Ser Val Asn Arg Ile Leu His Asp Leu Lys Asp Glu Lys Val Ile Gly
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Val Ile Asp Asn Lys Ile Met Ile Tyr Asn Pro Gln Tyr Leu Glu Glu
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Tyr Ile Ser
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Asp Thr Phe Leu Phe Gln Glu Gly Met Asp Ala Glu Glu Leu Tyr Leu
35 40 45

Ile Gln Ser Gly Leu Val Gln Ile Gly Lys Leu Thr Ser Asp Gly Lys
50 55 60

Glu Leu Thr Leu Arg Met Cys Lys Lys Asn Asp Ile Val Gly Glu Leu
65 70 75 80

Thr Leu Phe Thr Glu Asp Ala Lys Tyr Met Leu Ser Ala Lys Ile Leu

85

90

95

Ser Asp Gly Glu Val Leu Val Ile Asn Lys Asp Lys Leu Glu Lys Glu
100 105 110

Leu Ile Gln Asn Gly Ala Leu Thr Phe Glu Phe Met Lys Trp Met Ser
115 120 125

Thr His Leu Arg Lys Ile Gln Ser Lys Ile Arg Asp Leu Leu Leu Asn
130 135 140

Gly Lys Lys Gly Ala Leu Tyr Ser Thr Leu Ile Arg Leu Ala Asn Ser
145 150 155 160

Tyr Gly Ile Thr Arg Ser Asp Gly Ile Leu Ile Asn Ile Val Leu Thr
165 170 175

Asn Gln Asp Leu Ala Lys Phe Cys Ala Ala Ala Arg Glu Ser Ile Asn
180 185 190

Arg Met Leu Ser Asp Leu Arg Lys Asn Gly Val Ile Ser Ile Glu Asp
195 200 205

Ser Gly Lys Ile Val Ile His Gln Ile Asn Tyr Leu Lys Arg Glu Ile
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Asp Cys Glu Asn Cys Pro Leu Glu Ile Cys Asn Ile Asp
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<210> 8
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